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sequence, subsequence display, backbone, repeat region

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WD Repeat example

Each repeat **sequence** is broken into columns that describe the structure of the ... The conserved portions of the WD **repeat region** itself were shadowed with X's ... The BLASTP search results are as follows. * The amino **subsequence** is too The model coordinates are for the **backbone** atoms and beta carbons only of ... bmerc-www.bu.edu/psa/wd-example.htm - Cached - Similar - ⌂ ⌂ [X]

The mutated **subsequence** problem and locating conserved genes ...

by HL Chan - 2005 - Cited by 6 - Related articles

(1999): A conserved gene rarely comprises the same entire **sequence** in the two hot spots where a number of mutations cluster together in the same **region**. **Backbone**. Find the MWCS of A and B. We call this **subsequence** as the ... We repeat such refinement until no more problematic pairs of intervals are left. ... bioinformatics.oxfordjournals.org/cgi/content/full/2/1/10/2271 - Similar - ⌂ ⌂ [X]

New Approximation Algorithms for Longest Common Subsequences ...

by L Bergroth - 1998 - Cited by 14 - Related articles

identical **sequence** of (not necessarily contiguous) symbols found in both strings.

There's no guarantee that points on the **backbone** belong ... which define the **region** to be studied, all symbols of Σ' length string after which the source begins to **repeat** the generated **sequence** is short, the dependencies ...

ieeexplore.ieee.org/iel4/5792/15455/00712980.pdf?arnumber=712980 - ⌂ ⌂ [X]

Key to acedb graphic display

Mar 14, 2001 ... Below is an image of a **sequence display** for the AGR database. ... of the page to see a zoomed in view of just that **subsequence**. You can then use the 'Zoom out.' button to see the **region** of **sequence** surrounding your gene of interest ... **Backbone** marker, Single position or Best of Multiple positions. ...

ukcrop.net/agr/sequence_display_key - Cached - Similar - ⌂ ⌂ [X]

Structural stability of short **subsequences** of the tropomyosin chain featuring a pseudo-repeating heptad of amino acids ... the near C-terminal(253-280 **subsequence**), a sub-**sequence**-particularly rich in potential intrachain clude that the contribution to the **backbone region** ... doi.wiley.com/10.1002/bip.360350113 - ⌂ ⌂ [X]

Site-Specific Phosphorylation of Lys-Ser-Pro Repeat Peptides from ...

by P Sharma - 1998 - Cited by 20 - Related articles - All 5 versions

Peptides 1 and 2 were derived from the NF-H **sequence** of mouse and human species, respectively, while peptides 3 and 4 were from the KSP **repeat region** of mouse ... and the listed **sequence**, mult is the number of additional times a **subsequence** occurs Figure 5 (a, top) Overlay of the **backbone** atoms of the 24 best ...

pubs.acs.org/cgi-bin/jtext?bichaw/37/14/abs/bi0727460 - ⌂ ⌂ [X]

Mauve: Multiple Alignment of Conserved Genomic **Sequence** With ...

by ACE Darling - 2004 - Cited by 210 - Related articles - All 21 versions

... been developed to **display** the rearrangement structure of several genome sequences. Anchoring its alignment in unique **subsequences** provides Mauve with ... When scored

only on **regions** considered **backbone sequence**, the accuracy is **Repeat analysis** using tools such as **Repeat Masker**, REPuter (Kurtz et al. genome.cs.hip.org/content/14/7/1394.full -

Conservation of genetic information

by LF Harris - Cited by 10 - Related articles - All 11 versions

The **regions** of nucleotide **subsequence similarity** to the GRE in the GR DBD occur

Therefore, we proposed that the zinc finger may serve as a structural **backbone** for ... critical regulatory element within the MMTV 5' long terminal **repeat**, ... **regions** which showed **genetic sequence similarity** to the GR DBD exons 3, ...

www.msi.umn.edu/Projects/mg90601/pnas1.html - Cached - Similar -

Comprehensive **sequence** analysis of the 182 predicted open reading ... [PubMed] Holm L, Sander C. Database algorithm for generating protein **backbone** and side-chain ... Intragenic **repeat** structure and identification of a family of ... Identification of common molecular **subsequences**. J Mol Biol. ... [PubMed]; Sor F, Chéret G, Fabre F, Faye G, Fukuhara H. Sequence of the HMR region on ... www.ncbi.nlm.nih.gov/pmc/articles/PMC2142145/ -

Cell - The SV40 72 bp repeat preferentially potentiates ...

by C Kanduri - 2000 - Cited by 85 - Related articles

... in Simian Virus 40 (SV40) and various **sequence** mutants and insertion derivatives. ... Hence, stable curvature in this **region** of the VP1 gene is due to the five ... within a **region** of the helix **backbone** that is intrinsically curved, ... VRE(IFN α), tetrameric repA, and tetrameric GAAAGT (a **subsequence** of repB) or ...

www.cell.com/abstract/0092-8674(83)90470-1 - Cached -

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